



P021

A E

Test: DELPHI / DIY (circle one)

Universidad de La Laguna (Spain), 26 January 2022

Your initials:

- **Question 1:** Is the candidate gene a transcription factor or part of a transcription complex?

Gene	Transcription factor / part of transcription complex? (Y/N)
HIF1A	
SMAD3	
TRAF2	
CEBPA	Si Si

- **Question 2:** Is the candidate gene involved in an immune signalling pathway, i.e. a pathway that is immune cell-specific or a generic pathway known to operate in immune cells (e.g. the JAK-STAT, MAPK, NF-kappa B or TNF signalling pathways)?

You can also consider blood cell-specific pathways like leukaemias.

Gene	Annotate relevant pathways
HIF1A	
SMAD3	
TRAF2	
CEBPA	Si DMIL AML

- **Question 3:** Can you write down the known protein binding partners (if any) of each candidate gene?

Gene	Annotate relevant pathways
HIF1A	
SMAD3	
TRAF2	
CEBPA	Foxo1 KDM4Δ

Foxo1

- **Question 4:** Can you write down the diseases each candidate gene is associated with?

Gene	Annotate relevant pathways
HIF1A	
SMAD3	
TRAF2	
CEBPA	AML

AML

- **Question 5:** For each candidate gene, can you find a genetic phenotype that produces an abnormal immune phenotype or disease?

Gene	Annotate relevant pathways
HIF1A	
SMAD3	
TRAF2	
CEBPA	enlarged lymph nodes eosinophil count

- **Question 6:** Are any of the candidate genes expressed in a type of immune cell or in a tissue or organ where immune cells are known to reside (e.g. the bone marrow, thymus, spleen and lymph nodes) ?

Gene	Annotate relevant pathways
HIF1A	
SMAD3	
TRAF2	
CEBPA	bone marrow bone marrow lymph node

- **Question 7:** Have any of the candidate genes been studied in the context of macrophages?

Gene	PMID of paper
HIF1A	
SMAD3	
TRAF2	
CEBPA	23023392

34407972

System Usability Scale

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	Strongly disagree				Strongly agree	
1. I think that I would like to use this system frequently	1	2	3	4	5	3
2. I found the system unnecessarily complex	1	2	3	4	5	3
3. I thought the system was easy to use	1	2	3	4	5	4
4. I think that I would need the support of a technical person to be able to use this system	1	2	3	4	5	4
5. I found the various functions in this system were well integrated	1	2	3	4	5	4
6. I thought there was too much inconsistency in this system	1	2	3	4	5	3
7. I would imagine that most people would learn to use this system very quickly	1	2	3	4	5	4
8. I found the system very cumbersome to use	1	2	3	4	5	4
9. I felt very confident using the system	1	2	3	4	5	3
10. I needed to learn a lot of things before I could get going with this system	1	2	3	4	5	4

$$\begin{array}{r}
 3 \\
 \times 4 \\
 \hline
 12 \\
 \times 6 \\
 \hline
 24 \\
 \hline
 36
 \end{array}$$

$$\begin{array}{r}
 36 \\
 \times 2.5 \\
 \hline
 = 90
 \end{array}$$

Figure 8.6

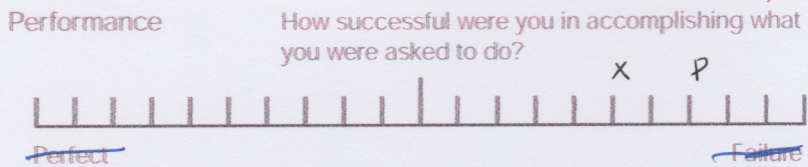
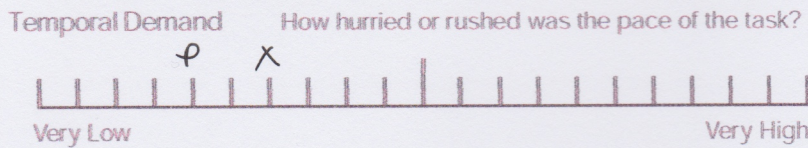
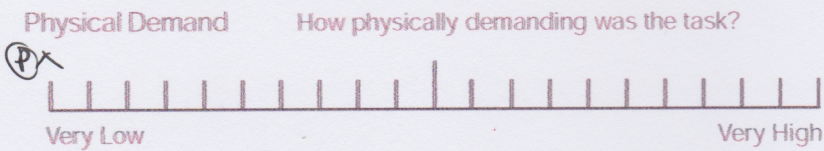
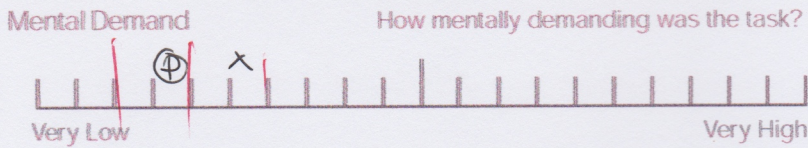
NASA Task Load Index

Hart and Staveland's NASA Task Load Index (TLX) method assesses work load on five 7-point scales. Increments of high, medium and low estimates for each point result in 21 gradations on the scales.

(P)

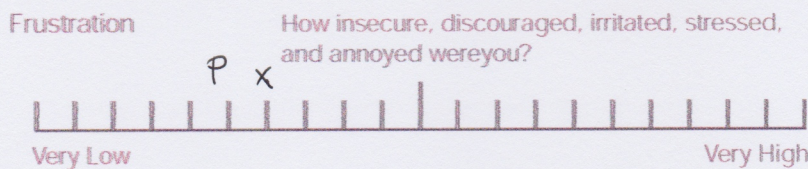
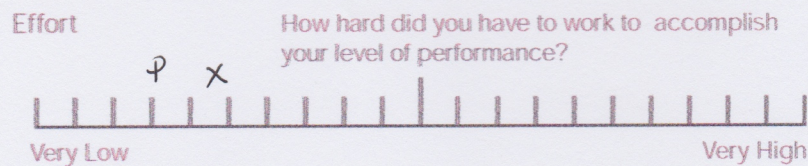
X

Name	Task	Date
------	------	------



very low

very high



PO21

~~XXXXXXXXXXXXXXXXXXXX~~

PhD student

~~XXXXXXXXXXXXXXXXXXXX~~

(P)

X

Q1

16"

PubMed

↓
NCBI

41"

Q2

16"

25"

Q3

41"

STRING-DB

1'02"

Q4

16"

28"

Q5

27"

NCBI gene → GeneCards 2'46"

Q6

29"

GeneCards

45"

Q7

34"

NCBI (all albs)

3'34"

↓
PubMed

Qué cosas te gustaría más preguntar?

⊕ - muy sencilla de utilizar y que le
información estar muy clara. clara

Que cambiarían?

⊖ que se permitiera poner una lista de preguntas
en vez de uno solo. = verbo en
conjunto

Algo más?

← Sería interesante tener una sí, información
estadística ~~de~~ de los resultados,
para no tener que crearlos por nosotros,
sino por un push por ejemplo.

e.g. (cómo de abundante es la expresión de expr?)

... en una población